

# Kalman Filter From The Ground Up

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## ESSENTIAL BACKGROUND

### MEAN AND EXPECTED VALUE

Suppose we want to compare the heights of two high school basketball teams. The following table provides the players' heights and the mean height of each team.

```
teamA <- c(1.89, 2.1, 1.75, 1.98, 1.85)
meanTeamA <- mean(teamA)
meanTeamA
```

```
[1] 1.914
```

```
teamB <- c(1.94, 1.9, 1.97, 1.89, 1.87)
meanTeamB <- mean(teamB)
meanTeamB
```

```
[1] 1.914
```

The distance from the mean for each variable would be

```
distanceFromMeanTeamA <- teamA - meanTeamA
distanceFromMeanTeamB <- teamB - meanTeamB
```

let's avoid negative numbers by squaring the distances

```
squaredDistanceFromMeanTeamA <- distanceFromMeanTeamA^2
squaredDistanceFromMeanTeamB <- distanceFromMeanTeamB^2
```

Sampled variance is the sum of squared distances from the mean divided by the number of observations minus one.

```
sampldVarianceTeamA <- sum(squaredDistanceFromMeanTeamA) / (length(teamA) - 1)
sampldVarianceTeamA
```

```
[1] 0.01763
```

```
sampldVarianceTeamB <- sum(squaredDistanceFromMeanTeamB) / (length(teamB) - 1)
sampldVarianceTeamB
```

```
[1] 0.00163
```

The variance is the average of the squared distances from the mean

```
varianceTeamA <- mean(squaredDistanceFromMeanTeamA)
varianceTeamA
```

```
[1] 0.014104
```

```
varianceTeamB <- mean(squaredDistanceFromMeanTeamB)
varianceTeamB
```

```
[1] 0.001304
```

The standard deviation is the square root of the variance

```
standardDeviationTeamA <- sqrt(varianceTeamA)
standardDeviationTeamA
```

```
[1] 0.1187603
```

```
standardDeviationTeamB <- sqrt(varianceTeamB)
standardDeviationTeamB
```

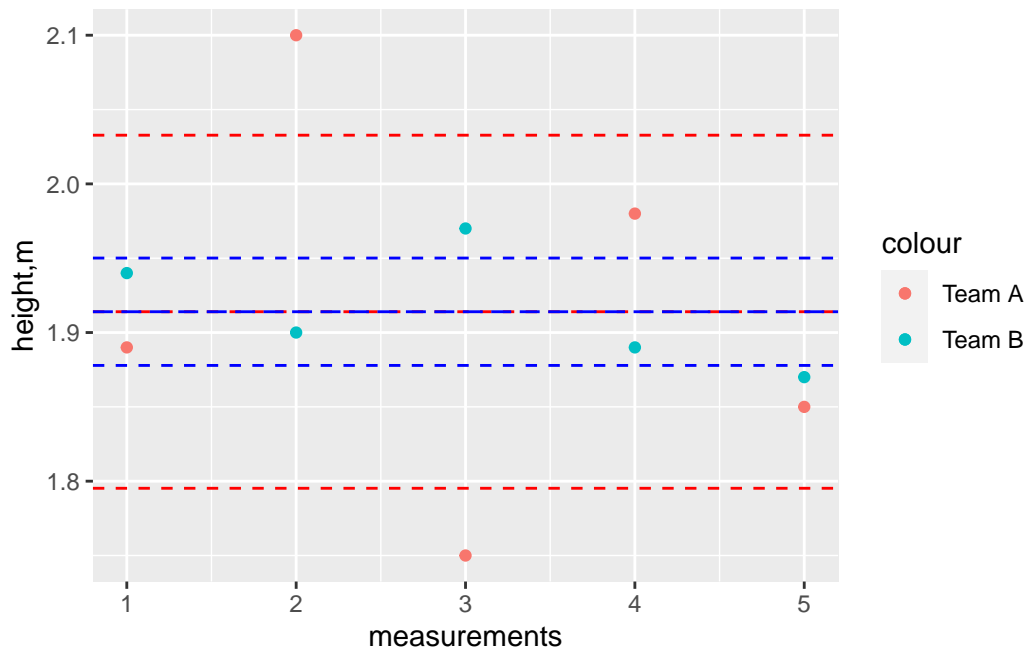
```
[1] 0.03611094
```

Final plot

```

ggplot() +
  labs(x = "measurements", y= "height,m") +
  geom_point(data = data.frame(teamA),
    aes(x = 1:length(teamA), y = teamA, color = "Team A")) +
  geom_point(data = data.frame(teamB),
    aes(x = 1:length(teamB), y = teamB, color = "Team B")) +
  geom_hline(yintercept = meanTeamA,
    color = "red", linetype = "dashed") +
  geom_hline(yintercept = meanTeamB,
    color = "blue", linetype = "longdash") +
  geom_hline(yintercept = meanTeamA + standardDeviationTeamA,
    color = "red", linetype = "dashed") +
  geom_hline(yintercept = meanTeamA - standardDeviationTeamA,
    color = "red", linetype = "dashed") +
  geom_hline(yintercept = meanTeamB + standardDeviationTeamB,
    color = "blue", linetype = "dashed") +
  geom_hline(yintercept = meanTeamB - standardDeviationTeamB,
    color = "blue", linetype = "dashed")

```



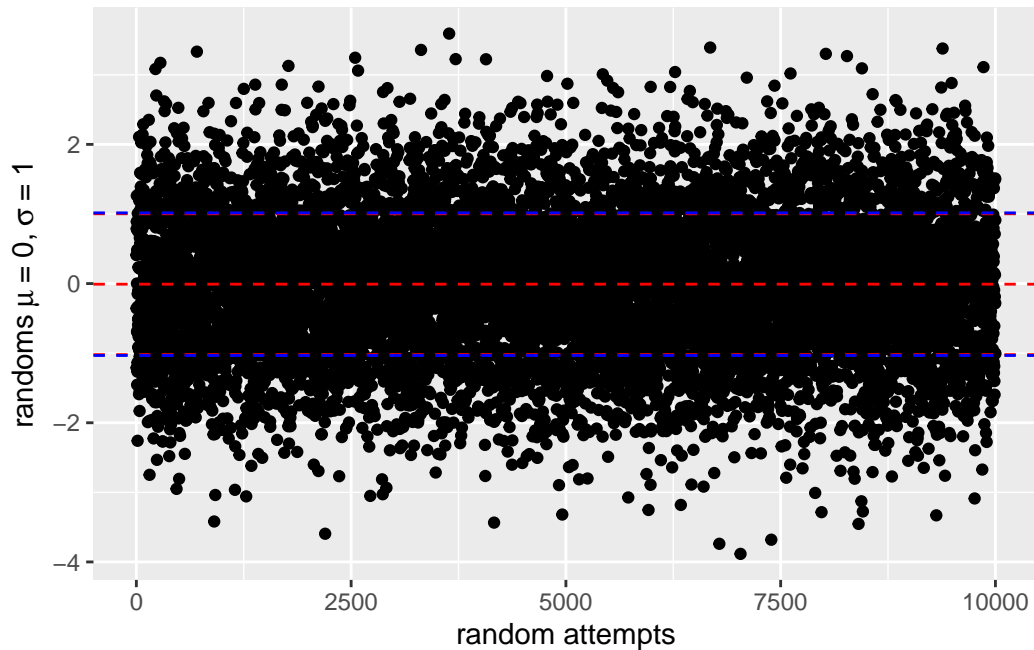
## NORMAL DISTRIBUTION

Generate a normal distribution with mean 0 and standard deviation 1

```

data <- rnorm(10000, mean = 0, sd = 1)
#, = 0, = 1
ggplot() +
  labs(x = "random attempts", y= expression("randoms" ~ mu ~ "=" ~ 0, " ~ sigma ~ "=" ~ 1)) +
  geom_point(data = data.frame(data),
             aes(x = 1:length(data), y = data)) +
  geom_hline(yintercept = mean(data),
             color = "red", linetype = "dashed") +
  geom_hline(yintercept = mean(data) + sd(data),
             color = "red", linetype = "dashed") +
  geom_hline(yintercept = mean(data) - sd(data),
             color = "red", linetype = "dashed") +
  geom_hline(yintercept = mean(data) + var(data),
             color = "blue", linetype = "dashed") +
  geom_hline(yintercept = mean(data) - var(data),
             color = "blue", linetype = "dashed")

```

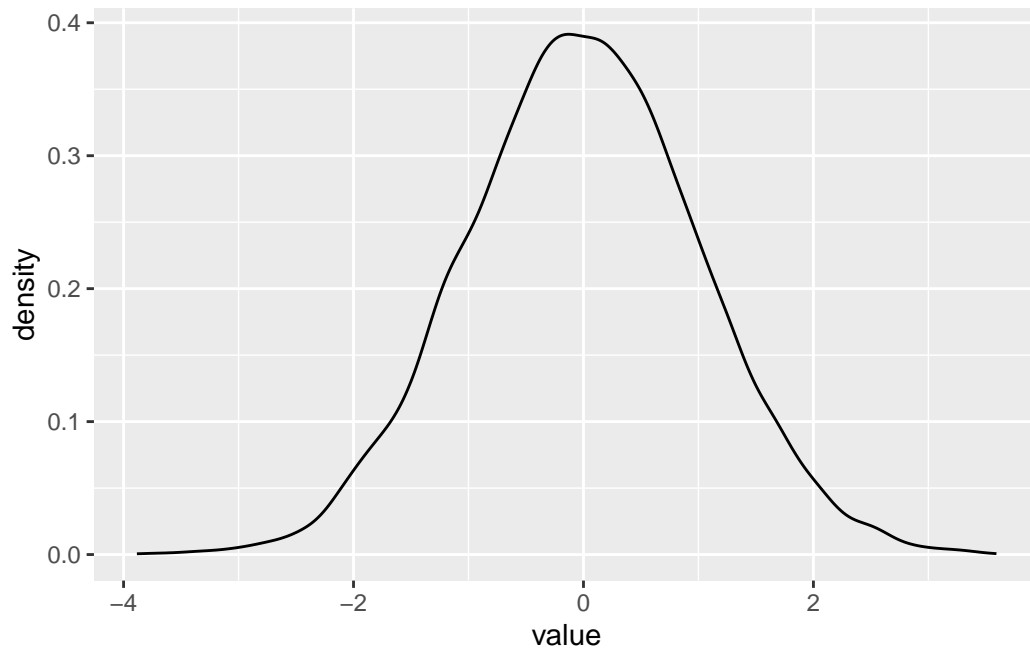


Plot distribution density with library function `geom_density`

```

ggplot() +
  labs(x = "value", y= "density") +
  geom_density(data = data.frame(data), aes(x = data))

```

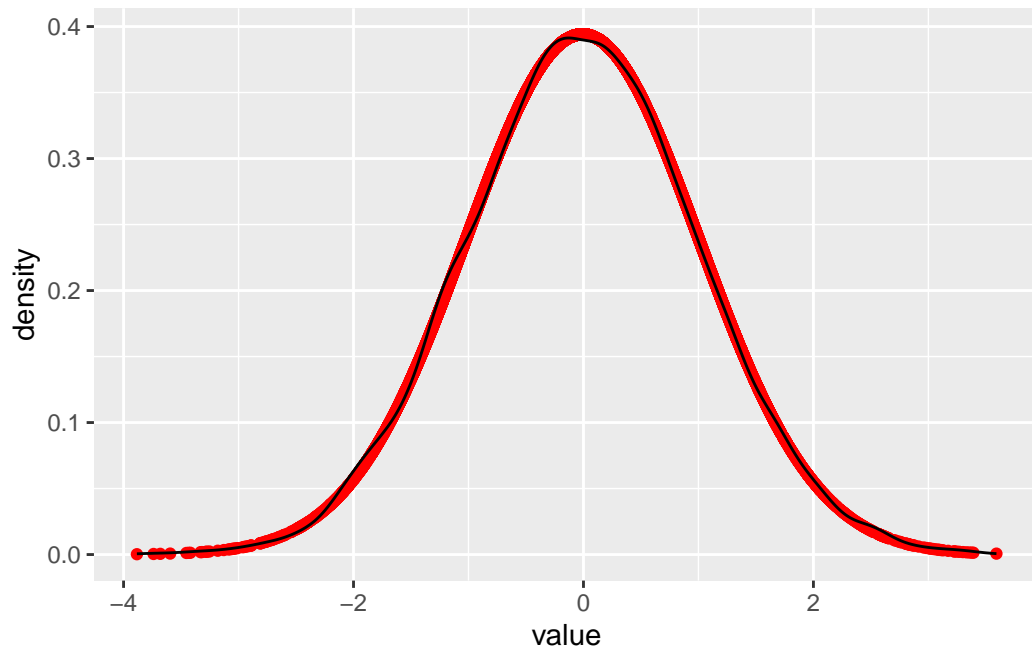


Calculated distribution density manually

```
mean <- mean(data)
variance <- var(data)
sdeviation <- sd(data)

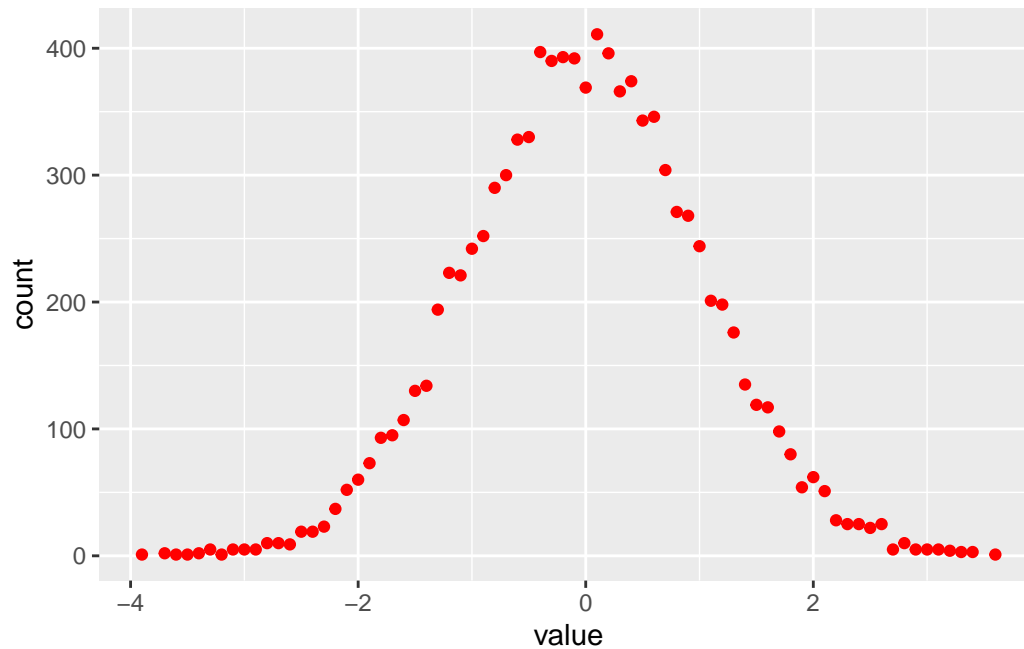
density <- function(x) {
  1 / sqrt(2 * pi * variance) * exp(-1/2 * (x - mean)^2 / variance)
}

dencityData <- dencity(data)
ggplot() +
  labs(x = "value", y = "density") +
  geom_point(data = data.frame(data, dencityData),
            aes(x = data, y = dencityData), color = "red")+
  geom_density(data = data.frame(data),
            aes(x = data))
```



Map each data to count of data rounded to 2 decimal places

```
dataCount <- data.frame(data) %>%  
  mutate(data = round(data, 1)) %>%  
  group_by(data) %>%  
  summarise(count = n())  
  
ggplot() +  
  labs(x = "value", y = "count") +  
  geom_point(data = dataCount, aes(x = data, y = count), color = "red")
```



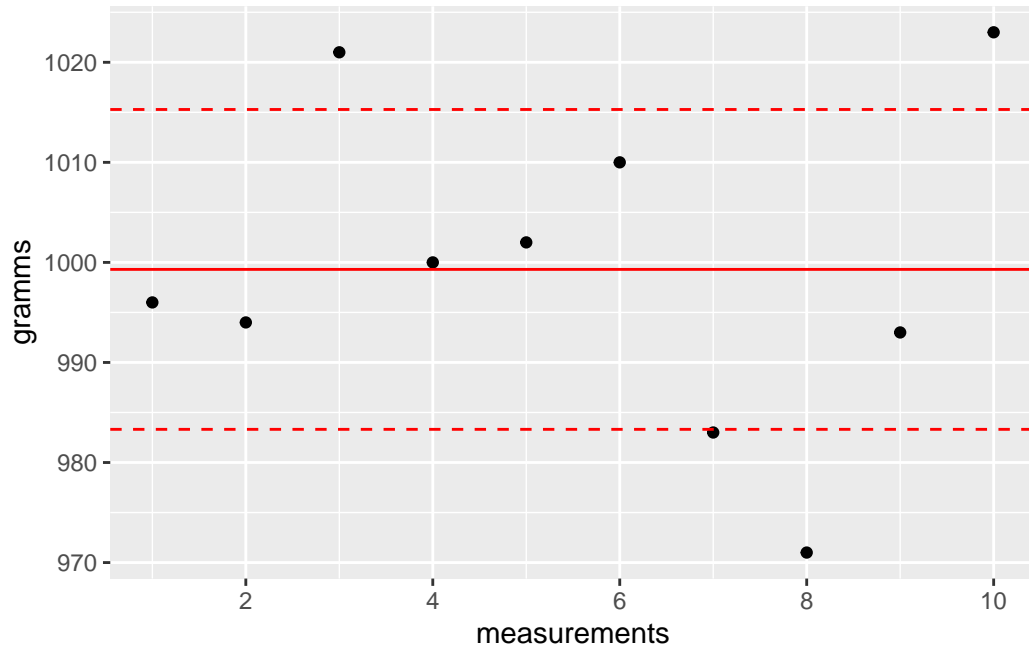
## THE $\alpha - \beta - \gamma$ FILTER

### Weighting the Gold Price

Collection of measurements

```
measurements <- c(996, 994, 1021, 1000, 1002, 1010, 983, 971, 993, 1000)

ggplot() +
  labs(x = "measurements", y = "grams") +
  scale_x_continuous( breaks=pretty_breaks()) +
  geom_point(data = data.frame(measurements),
             aes(x = 1:length(measurements), y = measurements)) +
  geom_hline(yintercept = mean(measurements),
             color = "red", linetype ="solid") +
  geom_hline(yintercept = mean(measurements) + sd(measurements),
             color = "red", linetype = "dashed") +
  geom_hline(yintercept = mean(measurements) - sd(measurements),
             color = "red", linetype = "dashed")
```



*Dynamic model of the system*

$$x_{n+1} = x_n \quad (1)$$

```
#' @description dynamic model of the system
# '
# ' @param x_n current state
# ' @returns predicted state

dynamicModel <- function(x_n) {
  return(x_n)
}
```

*State Update Equation*

$$x_n = x_{n-1} + \alpha(z_n - x_{n-1}) \quad (2)$$

where  $\alpha = \frac{1}{n}$

```
#' @description State Update Equation
# '
# ' @param x_n_1 previous estimation
```



```

#' @param z_n current measurement
#' @returns current updated estimation

stateEquation <- function(n, x_n_1, z_n) {
  alpha_n = 1/n
  innovation = (z_n - x_n_1)
  return(x_n_1 + alpha_n*innovation)
}

```

The Filter ( $\alpha$ ,  $\beta$  often are replaced with  $g$  and  $h$ )

```

#' @description The Filter
#'
#' @param x_0 initial state
#' @param Z measurements
#' @returns array of calculated states
filterG <- function(x_0, Z) {
  # predict
  results = c()
  x_n_1 = dynamicModel(x_0)
  n = 1
  for (z_n in measurements) {
    # estimate
    x_n = stateEquation(n, x_n_1, z_n)
    # add to estimations
    results = c(results, x_n)
    # update
    x_n_1 = dynamicModel(x_n)
    n = n + 1
  }
  return(results)
}

```

Apply the filter with the initial state of the first measurement

```

estimations <- filterG(measurements[c(1)], measurements)

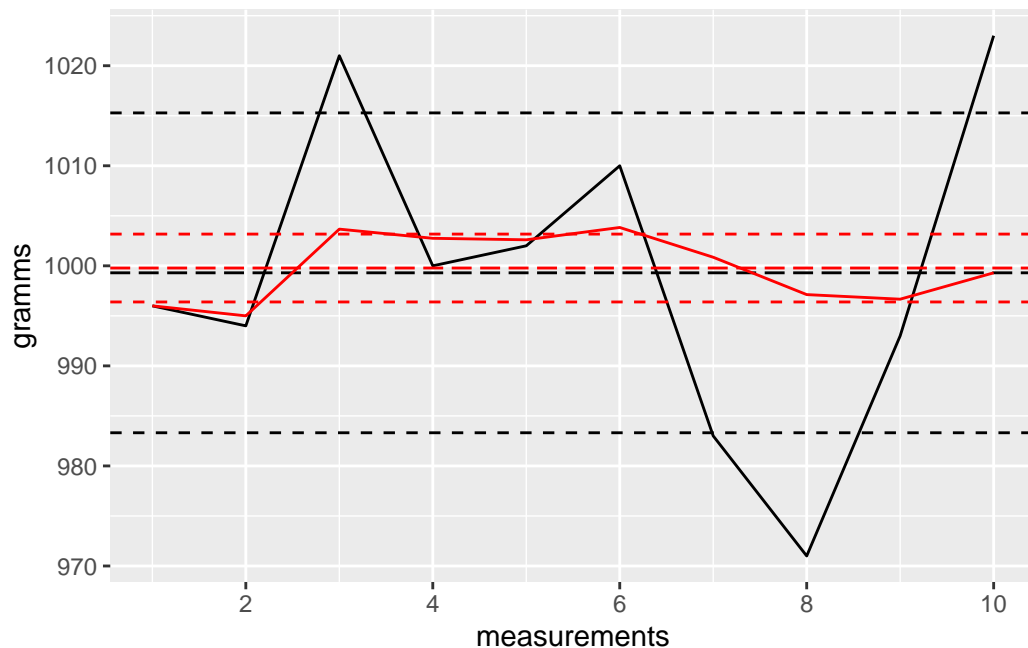
ggplot() +
  labs(x = "measurements", y= "gramms") +
  scale_x_continuous( breaks=pretty_breaks()) +
  geom_line(data = data.frame(measurements),
            aes(x = 1:length(measurements), y = measurements)) +

```

```

geom_hline(yintercept = mean(measurements),
           linetype = "longdash") +
geom_hline(yintercept = mean(measurements) + sd(measurements),
           linetype = "dashed") +
geom_hline(yintercept = mean(measurements) - sd(measurements),
           linetype = "dashed") +
geom_line(data = data.frame(estimations),
          aes(x = 1:length(estimations), y = estimations), color = "red")+
geom_hline(yintercept = mean(estimations),
           color = "red", linetype = "longdash") +
geom_hline(yintercept = mean(estimations) + sd(estimations),
           color = "red", linetype = "dashed") +
geom_hline(yintercept = mean(estimations) - sd(estimations),
           color = "red", linetype = "dashed")

```



### Constant Velocity

We assume an aircraft that is moving radially away from the radar (or towards the radar).

Collection of measurements

```

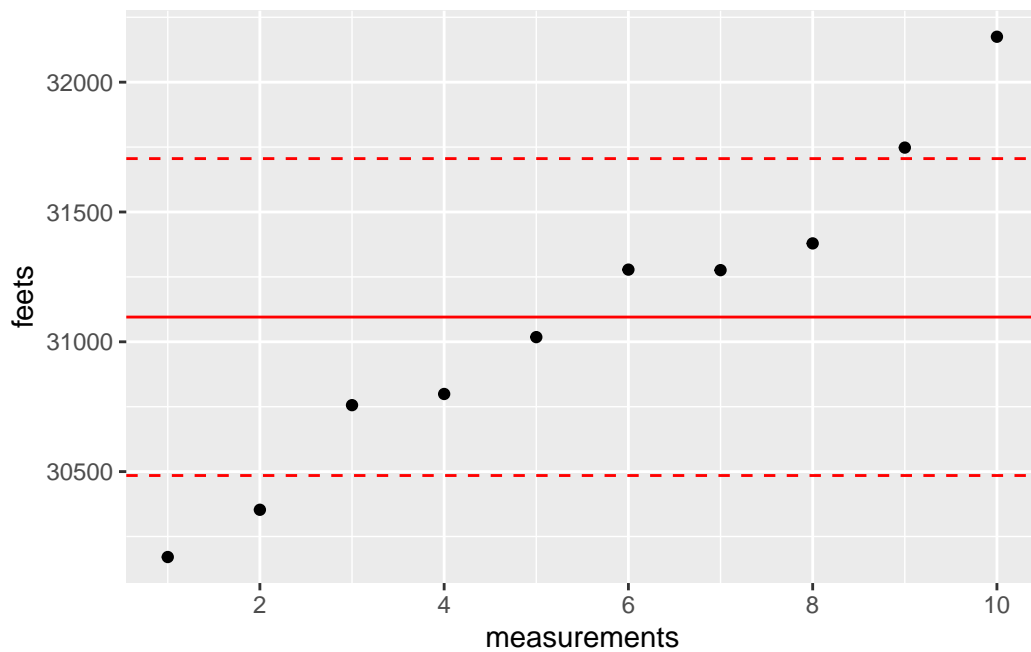
measurements <- c(30171, 30353, 30756, 30799, 31018, 31278, 31276, 31379,
                 31748, 32175)

ggplot() +
  labs(x = "measurements", y = "feets") +
  scale_x_continuous( breaks=pretty_breaks()) +
  scale_x_continuous(breaks = pretty_breaks()) +
  geom_point(data = data.frame(measurements),
            aes(x = 1:length(measurements), y = measurements)) +
  geom_hline(yintercept = mean(measurements),
            color = "red", linetype = "solid") +
  geom_hline(yintercept = mean(measurements) + sd(measurements),
            color = "red", linetype = "dashed") +
  geom_hline(yintercept = mean(measurements) - sd(measurements),
            color = "red", linetype = "dashed")

```

Scale for x is already present.

Adding another scale for x, which will replace the existing scale.



If  $x$  - is range, then velocity  $\dot{x}$  can be a derivative of the range :

$$\dot{x} = v = \frac{dx}{dt} \text{endequation} \quad (3)$$

*Dynamic model* (also called *State Extrapolation Equation* or *Transition Equation* or a *Prediction Equation*):

$$\begin{cases} x_{n+1} = x_n + \Delta t \dot{x}_n \\ \dot{x}_{n+1} = \dot{x}_n \end{cases} \quad (4)$$

In the matrix form:

$$\begin{pmatrix} x \\ \dot{x} \end{pmatrix} = \begin{pmatrix} 1 & \Delta t \\ 0 & 1 \end{pmatrix} \begin{pmatrix} x_n \\ \dot{x}_n \end{pmatrix} \quad (5)$$

```
#' @description dynamic model of the system
#'
#' @param dt time interval
#' @param x_n vector of previous state
#' @returns current state

dynamicModel <- function(dt, x_n) {
  A = matrix(c(1, dt, 0, 1), 2,2, TRUE)
  return(A %*% x_n)
}
```

Let's  $z_n$  be a measurement at the  $\Delta t$  time interval. Then the state update equation for velocity can be written as:

$$\hat{\dot{x}} = \hat{\dot{x}}_{n-1} - \beta \left( \frac{z_n - \hat{x}_{n-1}}{\Delta t} \right) \quad (6)$$

where  $\hat{\dot{x}}$  is the estimated state and  $\beta$  is the weighting factor to scale the impact of the new measurement.

And the *state update equation* for position can be written as:

$$\hat{x} = \hat{x}_{n-1} + \alpha (z_n - \hat{x}_{n-1}) \quad (7)$$

where  $\alpha$  is the weighting factor to scale the impact of the new measurement.

where  $\alpha$  is the weighting factor to scale the impact of the new measurement.

Unlike the “Weighting the Gold Price” example  $\alpha$  is constant and depends on the measurement precision.

```
#' @description State Update Equation
#
#' @param dt time interval needed to derive velocity
#' @param x_n_1 previous estimation vector
#' @param z_n current measurement
#' @returns current updated estimation

stateEquation <- function(alpha, beta, dt, x_n_1, z_n) {
  xHat_n_1 = x_n_1[1,1]
  innovationX = (z_n - xHat_n_1)
  xHat_n = xHat_n_1 + alpha*innovationX

  vHat_n_1 = x_n_1[2,1]
  innovationV = (z_n - xHat_n_1)/dt
  vHat = (vHat_n_1 + beta*innovationV)

  return(matrix(c(xHat_n, vHat), nrow = 2, ncol = 1))
}
```

The filter

```
#' @description The Filter
#
#' @param x_0 initial state
#' @param Z measurements
#' @returns array of calculated states

filterGF <- function(alpha, beta, x_0, Z, dt) {
  # setup time interval between measurements 5 seconds
  # predict
  results = list()
  x_n_1 = dynamicModel(dt, x_0)
  for (z_n in Z) {
    # estimate
    x_n = stateEquation(alpha, beta, dt, x_n_1, z_n)
    # add to estimations
    results = c(results, list(x_n))
    # update
    x_n_1 = dynamicModel(dt, x_n)
  }
}
```

```

    return(results)
  }

```

Apply the filter with the initial state of the first measurement with  $\alpha = 0.2$  and  $\beta = 0.1$

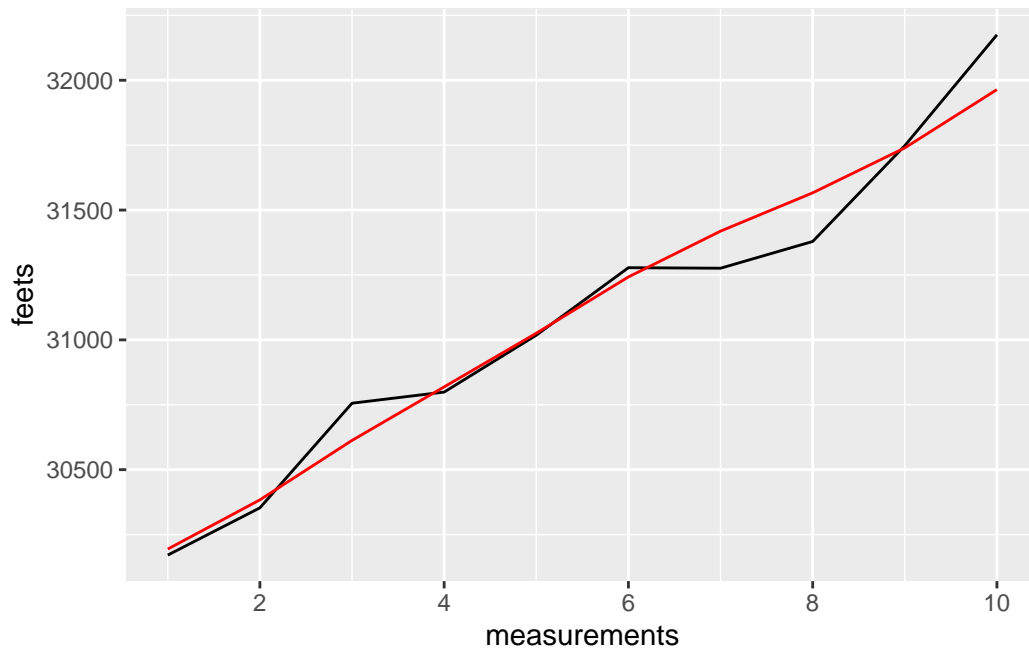
```

estimations <- filterGF(0.2, 0.1, matrix(c(30000, 40),2), measurements,5)

estimationsX <- unlist(sapply(estimations, function(x) x[1]))
estimationsV <- unlist(sapply(estimations, function(x) x[2]))

ggplot() +
  labs(x = "measurements", y= "feets") +
  scale_x_continuous( breaks=pretty_breaks()) +
  geom_line(data = data.frame(measurements),
            aes(x = 1:length(measurements), y = measurements)) +
  geom_line(data = data.frame(estimationsX),
            aes(x = 1:length(estimationsX), y = estimationsX), color = "red")

```



Apply the filter with the initial state of the first measurement with  $\alpha = 0.8$  and  $\beta = 0.5$

```

estimations <- filterGF(0.8, 0.5, matrix(c(30000, 40),2), measurements,5)

```

```

estimationsX <- unlist(sapply(estimations, function(x) x[1]))
estimationsV <- unlist(sapply(estimations, function(x) x[2]))

ggplot() +
  labs(x = "measurements", y = "feets") +
  scale_x_continuous( breaks=pretty_breaks()) +
  geom_line(data = data.frame(measurements),
            aes(x = 1:length(measurements), y = measurements)) +
  geom_line(data = data.frame(estimationsX),
            aes(x = 1:length(estimationsX), y = estimationsX), color = "red")

```

